DESIGN OF CDR-GRAFTED ANTI-RSV F PROTEIN VH

Glu Val Gin Leu Gin Gin Ser Gly Ala Glu Leu Val Arg Pro Gly Ala Leu Val Lys Leu Murine 1308F VH 5 10 15 20 Gln Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala Ser Val Lys Val Human HV3 VH Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala Ser Val Lys Val "CDR Grafted" VH

Gys Lys Ala Ser Gly Has Asn Ile Lys han Tur Tur Ile Tur Trp Val Arg Gln Ala 25 30 35 40 Ser Cys Lys Ala Ser Gly Tyr Thr Phe Asn Ser Tyr Tyr Met His Trp Val Arg Gln Ala lys lys Ala Sax Cly Phe Asn Ile lys Asp Tyz Tyz Ile Tyz Tzp Val lys Cln Ary

Pro Gly Gin Gly Leu Glu Tro Ile Gly Tro Ile Asp Pro Glu Asn Gly Asn The Val Phe OR 2
Pro Glu Gin Gly Leu Glu Tro Ile Gly Tro Ile Asp Pro Glu Asn Gly Asn The Val Phe 45 50 55 Fro Gly Gln Gly Leu Glu Trp Het Gly Ile Ile Asn Pro Ser Gly Gly Ser Thr Ser Tyr

65 70 75 80 Ala Gin Lys Phe Gin Gly Arg Val Thr Met Thr Arg Asp Thr Ser Thr Ser Thr Val Tyr hap Bro Lys Fins Gin Gly Ary Val Thr Met Thr Ary Asp Thr Ser Thr Ser Thr Val Tyr len I'm lyn I'hn Gin Gly lys Ala Ser lle Thr Ser Asp Thr Ser Ser Asn Thr Ala Tyr

Less Gin Less Sear Sear Less Thar Sear Gilu Asp Thar Ala Val Tyr Tyr Cys Ala Tyr Tyr Gly Lou Ser Ser Lou Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys Ala Tyr Tyr Gly 95 Yu Lou Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys Ala 18

The Sec Sec The Asp The Trp Gly Gln Gly The The Leu The Val Sec Sec Our 3

The Sec Sec The Asp The Trp Gly Gln Gly The The Leu The Val Sec Sec Inc. (1988)

DESIGN OF CDR-GRAFTED ANTI-RSV F PROTEIN VL

Ile Thr Cys Iva Ala Ser Gin Asp Ile Asn Ard Tyr Len Asn Trp Tyr Gin Gin Iys Pro Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Ser Trp Leu Ala Trp Tyr Gln Gln Lys Pro Asp II.e Lys Met The Gin See Pro See See Met Tyr Val See Leu Gly Glu Arg Val The - Marine 1308F VI 5 10 15 20 Asp Ile Gln Met Thr Gln Ser Pro Ser Thr Leu Ser Ala Ser Val Gly Asp Arg Val Thr Asp II Gin Met Thr Gin Ser Pro Ser Thr Leu Ser Ala Ser Val Gly Asp Arg Val Thr - "ODR Grafted" VL - Human K102 VI

Gly Lye Ser Pro Lys The Lou Lie His Ary Ala Asn Ary Lou Val Asp Gly Val Pro Ser Gly Lys Ala Pro Lys Leu Leu Ile Tyr Arg Ala Asp Arg Leu Val Asp Gly Val Pro Ser Gly Lys Ala Pro Lys Leu Leu Ile Tyr Asp Ala Ser Ser Leu Glu Ser Gly Val Pro Ser

He the Oys Lys Ala Ser Gin Asp He Asn Ary Tyr Leu Asn Trp Fhe Gin Gin Lys Pro

65 70 75 80 Arg Phe Ser Gly Ser Gly Ser Gly Thr Glu Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro Any Pho See Cly See Cly See Cly Cln Clu Tyr See Leu The Ile See See Leu Clu Pho Ary Phe Ser Cly Ser Cly Ser Cly Thr Clu Phe Thr Leu Thr Ile Ser Ser Leu Cln Pro

Asp Asp Fine Ala The Tyr Tyr Cys Len Gin Pha His Giu Pha Pro Tyr The Phe Gly Gly 85 90 95
Asp Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Tyr Asn Ser Tyr Ser Cin Asp Het Cily Lie Tyr Tyr Cys Leu Cin Phe His Ciu Phe Pro Tyr Thr Phe Cily Cily 100

Gly The Lys Leu Glu Ile Lys Gly The Lys Leu Glu Ile Lys



Figure 3 oligie used to make Hul308 VH

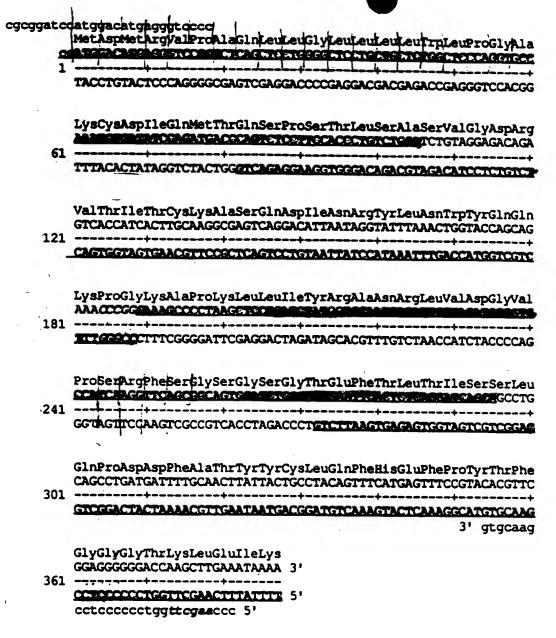


Figure 4. Oligo's used to make H2308 VL

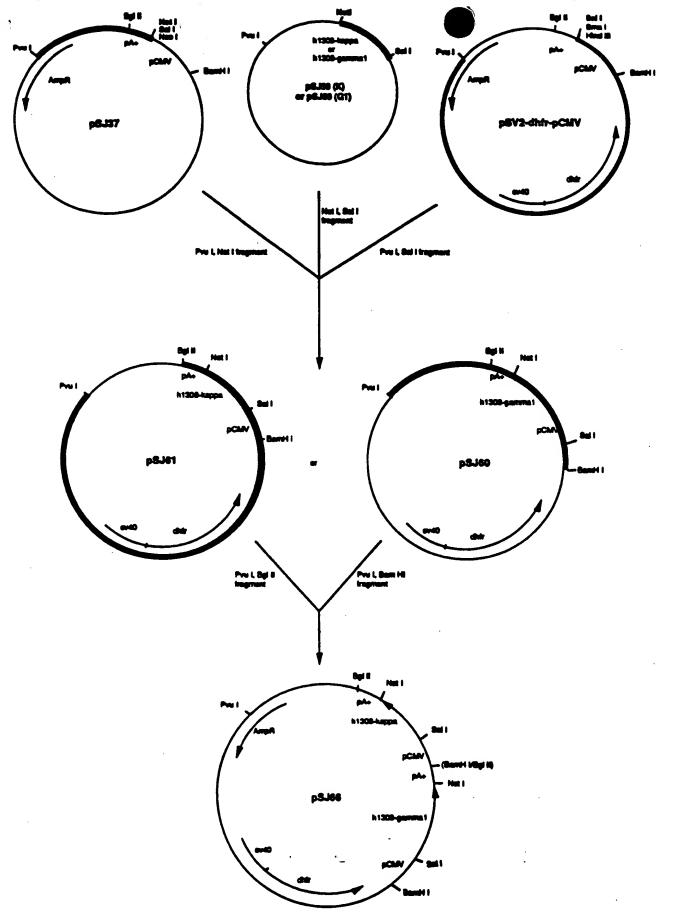


Fig 5. Construction of the Humized 1308 expression rachers

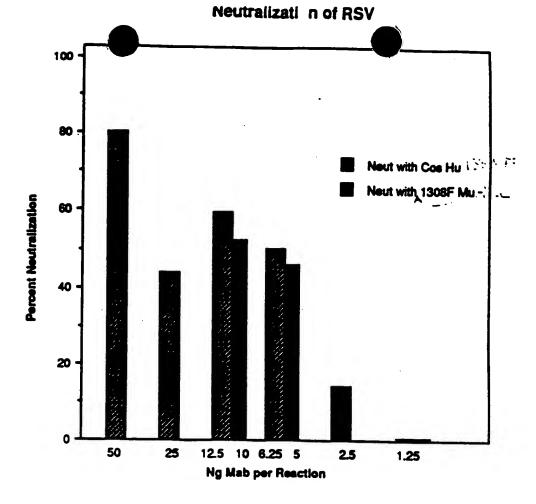


FIGURE 6

Design of Humanized VH f r anti-RSV Mab 1129

Gln Val Thr Leu Arg Glu Ser Gly Pro Ala Leu Val Lys Pro Ser Human VH (Cor)

1 Gln Val Thr Leu Arg Glu Ser Gly Pro Ala Leu Val Lys Pro (Ser) "Humanized" VH
Gln Val Glu Leu Gln Glu Ser Gly Pro Gly Ile Leu Gln Pro Ser Murine 1129 VH

Gln Thr Leu Thr Leu Thr Cys Ser Phe Ser Gly Phe Ser Leu Ser 16 Gln Thr Leu Thr Leu Thr Cys Ser Phe Ser Gly Phe Ser Leu Ser Gln Thr Leu Ser Leu Thr Cys Ser Phe Ser Gly Phe Ser Leu Ser

Ser Ser Gly Met Cys Val Gly Trp Ile Arg Gln Pro Pro Gly Lys

31 Ihr Ser Gly Met Ser Val Gly Trp Ile Arg Gln Pro Pro Gly Lys
Thr Ser Gly Met Ser Val Gly Trp Ile Arg Gln Pro Ser Gly Glu

Ala Leu Glu Trp Leu Ala Asp Ile Glu Trp Asp Asp Lys Asp
46 Alu Leu Glu Trp Leu Ala Asp Ile Trp Trp Asp Asp Lys Lys Asp
Gly Leu Glu Trp Leu Ala Asp Ile Trp Trp Asp Asp Lys Lys Asp

Tyr Asn Thr Ser Leu Asp Thr Arg Leu Thr Ile Ser Lys Asp Thr
61 Tyr Asn Pro Ser Leu Lys Ser Arg Leu Thr Ile Ser Lys Asp Thr
Tyr Asn Pro Ser Leu Lys Ser Arg Leu Thr Ile Ser Lys Asp Thr

Ser Lys Asn Gln Val Val Leu <u>Vrhr Val Thr</u> Asn Val Asp Pro Ala 76 Ser Lys Asn Gln Val Val Leu <u>Lys</u> Val Thr Asn <u>Val</u> asp Pro Ala Ser Ser Asn Gln Val Phe Leu Lys Ile Thr Gly Val Asp Thr Ala

Asp Thr Alu Thr Tyr Tyr Cys Alu Arg Ile Thr Val Ile Pro Alu Pro Alu Gly
91 Asp Thr Alu Thr Tyr Tyr Cys Alu Arg Ser Met Ile Thr Asn Trp - - Asp Thr Alu Thr Tyr Tyr Cys Alu Arg Ser Met Ile Thr Asn Trp - - -

Tyr Met Asp Val Trp Gly Arg Gly Thr Pro Val Thr Val Ser Ser

106 Tyr Phe Asp Val Trp Gly Ala Gly Thr Thr Val Thr Val Ser Ser

Tyr Phe Asp Val Trp Gly Ala Gly Thr Thr Val Thr Val Ser Ser

Asp Ile Gln Met Thr Gln Ser Pro Ser Thr Leu Ser Ala Ser Val Asp Ile Gln Met Thr Gln Ser Pro Ser Thr Leu Ser Ala Ser Val Asp Ile Gln Leu Thr Gln Ser Pro Ala Ile Met Ser Ala Ser Pro

- Human K102 VL
- "CDR Grafted" VL
- Murine 1129 VL

Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Gly Asp Arg Val Thr Ile Thr Cys Lys Cys Gln Leu Ser Val Gly Gly Glu Lys Val Thr Met Thr Cys Ser Ala Ser Ser Val Gly Ser Trp Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Tyr Met flis - Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Tyr Met His - Trp Tyr Gln Gln Lys Ser Ser Thr Ser Pro Lys Leu Leu Ile Tyr Asp Ala Ser Ser Leu Glu Ser Gly Val Pro Ser Leu Trp Ile Tyr Asp Thr Ser Lys Leu Ala Ser Gly Val Pro Ser Leu Trp Ile Tyr Asp Thr Ser Lys Leu Ala Ser Gly Val Pro Gly Arg Phe Ser Gly Ser Gly Ser Gly Thr Glu Phe Thr Leu Thr Ile Arg Phe Ser Gly Ser Gly Ser Gly Thr Glu Phe Thr Leu Thr Ile Arg Phe Ser Gly Ser Gly Ser Gly Asn Ser Tyr Ser Leu Thr Ile Ser Ser Leu Gln Pro Asp Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Ser Leu Gln Pro Asp Asp Phe Ala Thr Tyr Tyr Cys Phe Gln Ser Ser Ile Gln Ala Glu Asp Val Ala Thr Tyr Tyr Cys Phe Gln 100 105

Tyr Asn Ser Tyr Ser

Gly Ser Gly Tyr Pro Phe Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys CDR 3 Gly Ser Gly Tyr Pro Phe Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys

<<V / J>>

Oligonucleotides for the construction of the Humanized 1129 VH gene segment

SJ150 5'-CCAG	
	60
MetAspTrpThrTrpArgValPreCysTexLetAlaValAlaProGlyAlaHisSerGin	-
GIVACCITAACEXAGIUTGGIUTGGGGGGGAAACCCACACACACCCTCACACICACC	
T'-CONSTIGRACITES	
ValThrLe.ArgGluSerGlyProAlaLeuVallysProThrGlnThrLeuThrLeuThr	-
TGCACC-3' 8J151 5'- CAG	100
· · · · · · · · · · · · · · · · · · ·	T80
CysTrrPheSerClyPheSerLeuSerThrSerGlyMetSerValGlyTrpIleArgGln	-
GGGGTCCCTTCCGCG-5' SJ149 3'- GATA	240
ProProG_yLysAlaLeuGluTrpLeuAlaAspIleTrpTrpAspAspLysLysAspTyr	
AATCCATCCCTGAAG-3' SJ152 5'-GGTC	
	300
AshProSerLeulysSerArgLouThrIleSerLysAspThrSerLysAshSlnVaiVal	•
CAATTTCACTCACTCG-5' SJ148 3'-TAC	360
LeuLySValThrAsnyM-ny-pProAlaAspltnAlaThrTyrTyrCysAlaArgSerMet	
	7
	MetAspTryThrTryArgValPerYslexLevAlaValAlaProGlyAlaHisSerGin GRACCITAAAXXAGICTGGTCTGCCCGGTCAAACCCCCACACCCCCACACCCTCACCCACC

